

trans-Bis(2-amino-1,3-benzothiazole- κN^3)-dichlorocopper(II)

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Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.011\text{ \AA}$

R factor = 0.038

wR factor = 0.087

Data-to-parameter ratio = 9.2

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title complex, $[\text{CuCl}_2(\text{C}_7\text{H}_6\text{N}_2\text{S})_2]$, contains a Cu centre with a distorted square-planar coordination geometry, involving two Cl ligands and two endocyclic N atoms from the thiazole moieties [$\text{Cu}-\text{Cl} = 2.275(2)$ and $2.297(2)\text{ \AA}$, and $\text{Cu}-\text{N} = 1.986(5)$ and $1.988(5)\text{ \AA}$]. The amino groups participate in intra- and intermolecular N–H···Cl hydrogen bonds, with N···Cl distances in the range $3.202(6)$ – $3.316(6)\text{ \AA}$.

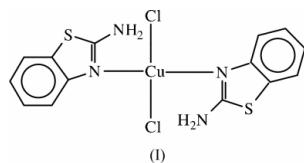
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Comment

Aminothiazolines are perceived as an important type of S,N-containing heterocycles (Raper, 1994). Usually, 2-substituted derivatives of 1,3-benzothiazole ($2-X\text{-btz}$; where X is NH_2 , CH_3 , Cl or S) act as σ -monodentate ligands through the ring N atom (Giusti & Peyronel, 1982). The coordination of 2-aminobenzothiazole (2-abtz) to the M^{II} ions of Co (Macíček *et al.*, 1987), Zn (Usman *et al.*, 2003) and Cu (Sieroń & Bukowska-Strzyżewska, 1999, 2000) has been studied thus far, and those studies confirm that heterocyclic nitrogen-bonded, rather than sulfur-bonded, complexes are found.



This paper reports the synthesis and crystal structure of a new copper(II) halide complex with 2-abtz, *trans*-bis(2-amino-1,3-benzothiazole- κN^3)-dichlorocopper(II), (I) (Fig. 1), which consists of discrete neutral molecules. These molecules are almost centrosymmetric. The copper centre is coordinated by two Cl ligands and two endocyclic N atoms from the thiazole

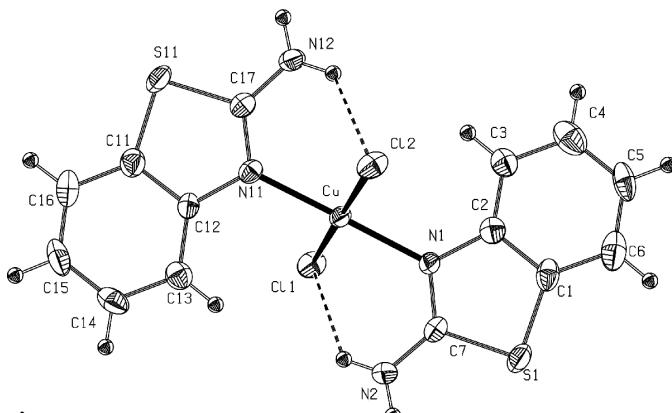


Figure 1

View of (I), with the atom-numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

moieties [$\text{Cu}-\text{Cl} = 2.275(2)$ and $2.297(2)$ Å, and $\text{Cu}-\text{N} = 1.986(5)$ and $1.988(5)$ Å]. The amino N atoms are not involved in the Cu coordination. Tetrahedral distortion of the square-planar coordination is observed with a tetrahedrality angle θ (Holm & O'Connor, 1971) of $2.6(2)^\circ$. The bond lengths in the 2-abtz ring system (Table 1) are normal (Allen *et al.*, 1987). Due to the pronounced delocalization in the $\text{S}-\text{C}\equiv\text{N}$ fragment of the thiazole ring, the $\text{S}1-\text{C}1$ [1.742(7) Å] and $\text{S}11-\text{C}11$ [1.733(7) Å] bonds are slightly shorter than $\text{S}1-\text{C}7$ [1.751(7) Å] and $\text{S}11-\text{C}17$ [1.752(6) Å]. Also, the bond angles around the endocyclic S atoms are in agreement with the literature values for the geometry around an endocyclic S atom [$\text{C}1-\text{S}1-\text{C}7 = 89.6(3)^\circ$ and $\text{C}11-\text{S}11-\text{C}17 = 89.9(3)^\circ$]. The ligands are planar, with the angles between the mean planes calculated through the five- and six-membered rings of each ligand being $2.4(3)$ and $3.4(3)^\circ$. The ligands are almost parallel, with an interplanar angle of $3.8(2)^\circ$. The amino groups are coplanar with the attached thiazole planes. In the crystal packing, each amino group is involved in intra- and intermolecular $\text{N}-\text{H}\cdots\text{Cl}$ hydrogen bonds (Table 2). The $\text{N}2-\text{H}2\cdots\text{Cl}1^i$ and $\text{N}12-\text{H}12\cdots\text{Cl}2^{ii}$ hydrogen bonds interconnect the molecules into chains along the **a** direction (Fig. 2).

Experimental

The title complex was prepared by addition of 0.001 mol of $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ to a warm solution containing 0.001 mol of ligand in 50 ml of ethanol. The resulting solution was filtered and allowed to cool. Slow evaporation deposited dark-brown crystals analyzed as $\text{CuCl}_2(\text{C}_7\text{H}_6\text{N}_2\text{S})_2$. Analysis found: C 38.20, N 13.02, H 2.93%; calculated: C 38.67, N 12.88, H 2.78%.

Crystal data

$[\text{CuCl}_2(\text{C}_7\text{H}_6\text{N}_2\text{S})_2]$	$D_m = 1.75(3)$ Mg m $^{-3}$
$M_r = 434.87$	D_m measured by flotation
Monoclinic, $P2_1$	Mo $\text{K}\alpha$ radiation
$a = 6.4430(10)$ Å	Cell parameters from 15 reflections
$b = 13.032(3)$ Å	$\theta = 10.0\text{--}15.0^\circ$
$c = 9.987(2)$ Å	$\mu = 1.96$ mm $^{-1}$
$\beta = 106.85(2)^\circ$	$T = 293$ K
$V = 802.6(3)$ Å 3	Needle, brown
$Z = 2$	$0.4 \times 0.1 \times 0.1$ mm
$D_x = 1.799$ Mg m $^{-3}$	0.4 × 0.1 × 0.1 mm

Data collection

Siemens <i>P3</i> diffractometer	$R_{\text{int}} = 0.026$
ω -2θ scans	$\theta_{\text{max}} = 27.6^\circ$
Absorption correction: ψ scan (North <i>et al.</i> , 1968)	$h = -8 \rightarrow 8$
$T_{\text{min}} = 0.721$, $T_{\text{max}} = 0.822$	$k = 0 \rightarrow 16$
2096 measured reflections	$l = 0 \rightarrow 12$
1933 independent reflections	3 standard reflections every 50 reflections
1649 reflections with $I > 2\sigma(I)$	intensity decay: none

Refinement

Refinement on F^2	$w = 1/[σ^2(F_o^2) + (0.0445P)^2 + 0.5431P]$
$R[F^2 > 2\sigma(F^2)] = 0.038$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.087$	$(Δ/\sigma)_{\text{max}} < 0.001$
$S = 1.03$	$Δρ_{\text{max}} = 0.96$ e Å $^{-3}$
1933 reflections	$Δρ_{\text{min}} = -0.44$ e Å $^{-3}$
209 parameters	Absolute structure: Flack (1983)
H-atom parameters constrained	Flack parameter = 0.08(3)

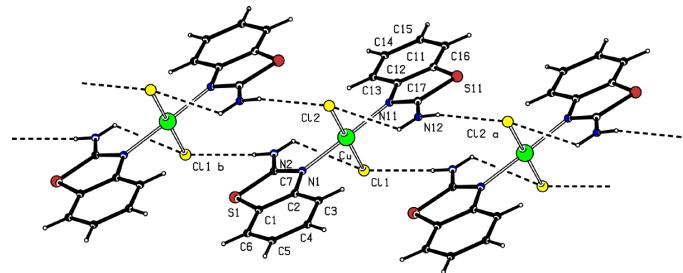


Figure 2

Diagram showing hydrogen bonds in (I) as dashed lines. [Symmetry codes: (a) $1+x, y, z$; (b) $x-1, y, z$, corresponding to (ii) and (i), respectively, in Table 2.]

Table 1

Selected geometric parameters (Å, °).

$\text{Cu}-\text{Cl}1$	2.2747(19)	$\text{Cu}-\text{N}1$	1.986(5)
$\text{Cu}-\text{Cl}2$	2.2969(19)	$\text{Cu}-\text{N}11$	1.988(5)
$\text{Cl}1-\text{Cu}-\text{Cl}2$	177.58(7)	$\text{Cl}2-\text{Cu}-\text{N}1$	89.98(19)
$\text{Cl}1-\text{Cu}-\text{N}1$	89.34(19)	$\text{Cl}2-\text{Cu}-\text{N}11$	89.41(19)
$\text{Cl}1-\text{Cu}-\text{N}11$	91.32(19)	$\text{N}1-\text{Cu}-\text{N}11$	178.7(2)

Table 2

Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N}2-\text{H}1\cdots\text{Cl}1$	0.86	2.74	3.266(6)	121
$\text{N}2-\text{H}2\cdots\text{Cl}1^i$	0.86	2.48	3.265(6)	152
$\text{N}12-\text{H}11\cdots\text{Cl}2$	0.86	2.64	3.202(6)	124
$\text{N}12-\text{H}12\cdots\text{Cl}2^{ii}$	0.86	2.54	3.316(6)	151

Symmetry codes: (i) $x-1, y, z$; (ii) $1+x, y, z$.

All H atoms were placed in calculated positions and refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$.

Data collection: *P3 Software* (Siemens, 1993); cell refinement: *P3 Software*; data reduction: *XDISK* in *SHELXTL/PC* (Sheldrick, 1990); structure solution: *SHELXS97* (Sheldrick, 1997); structure refinement: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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